

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 09/441,242B

Source: IFw/b

Date Processed by STIC: 7/14/06

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 07/14/2006

PATENT APPLICATION: US/09/441,242B

TIME: 09:20:14

Input Set : F:\3589.1017-001 Seq List.txt

Output Set: N:\CRF4\07142006\I441242B.raw

4 <110> APPLICANT: Russo, Giandomenico
 5 Croce, Carlo M.
 7 <120> TITLE OF INVENTION: TCL-1 GENE AND PROTEIN AND RELATED
 8 METHODS AND COMPOSITIONS
 10 <130> FILE REFERENCE: 3589.1017-001
 12 <140> CURRENT APPLICATION NUMBER: 09/441,242B
 13 <141> CURRENT FILING DATE: 1999-11-16
 15 <150> PRIOR APPLICATION NUMBER: 08/330,272
 16 <151> PRIOR FILING DATE: 1994-10-27
 18 <160> NUMBER OF SEQ ID NOS: 12
 20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 22 <210> SEQ ID NO: 1
 23 <211> LENGTH: 1324
 24 <212> TYPE: DNA
 25 <213> ORGANISM: Artificial Sequence
 27 <220> FEATURE:
 28 <223> OTHER INFORMATION: cDNA sequence of TCL-1
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 31 <222> LOCATION: (49)...(387)
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 35 Ala Glu Cys
 36 1
 38 ccg aca ctc ggg gag gca gtc acc gac cac ccg gac cgc ctg tgg gcc 105
 39 Pro Thr Leu Gly Glu Ala Val Thr Asp His Pro Asp Arg Leu Trp Ala
 40 5 10 15
 42 tgg gag aag ttc gtg tat ttg gac gag aag cag cac gcc tgg ctg ccc 153
 43 Trp Glu Lys Phe Val Tyr Leu Asp Glu Lys Gln His Ala Trp Leu Pro
 44 20 25 30 35
 46 tta acc atc gag ata aag gat agg tta cag tta cgg gtg ctc ttg cgt 201
 47 Leu Thr Ile Glu Ile Lys Asp Arg Leu Gln Leu Arg Val Leu Leu Arg
 48 40 45 50
 50 cgg gaa gac gtc gtc ctg ggg agg cct atg acc ccc acc cag ata ggc 249
 51 Arg Glu Asp Val Val Leu Gly Arg Pro Met Thr Pro Thr Gln Ile Gly
 52 55 60 65
 54 cca agc ctg ctg cct atc atg tgg cag ctc tac cct gat gga cga tac 297
 55 Pro Ser Leu Leu Pro Ile Met Trp Gln Leu Tyr Pro Asp Gly Arg Tyr
 56 70 75 80
 58 cga tcc tca gac tcc agt ttc tgg cgc tta gtg tac cac atc aag att 345
 59 Arg Ser Ser Asp Ser Ser Phe Trp Arg Leu Val Tyr His Ile Lys Ile
 60 85 90 95
 62 gac ggc gtg gag gac atg ctt ctc gag ctg ctg cca gat gac 387
 63 Asp Gly Val Glu Asp Met Leu Leu Glu Leu Leu Pro Asp Asp

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64 100                105                110
66 tgatgtatgg tcttggcagc acctgtctcc ttaccacca gggcctgagc ctggccagcc 447
67 tacaatgggg atgtgtgtgt tctgttcacc ttcgtttact agcctgtgtg ctctccacc 507
68 acgcgtggggg ctggggaggaa tggacagaca gaggatgagc tctaccaggg gacctgcagg 567
69 cctgcctgta gcccaactctg ctgcgccttag cactaccact cctgccaaag aggattccat 627
70 ttggcagagc ttcttccagg tgcccagcta tacctgtgcc tcggcttttc tcagctggat 687
71 gatggtcttc agcctctcttc tgtcccttct gtccctcaca gcactagat ttcattgtgc 747
72 acacccactc agctcctgta acctgtgaga acacagccga ttcacctgag caggacctct 807
73 gaaacccctgg accagtggtc tcacatgggt ctacgcctgc atgtaaacac gctgcgaaac 867
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75 tgctgcacca cacagggttca cgtgcagctc aaggaaaggg ctgaaaggag cccttatctc 987
76 tgctcaggac tcagaagcct ctgggtcagt ggtccacatc ccgggacgca gcaggaggcc 1047
77 aggcggcgga gccctgtgga tgagccctca gaacccctgg cttgccacgc tggaaaaagg 1107
78 atagaggttg ggtttccccc ctttatagat ggtcacgcac ctgggttgta caaagtgtga 1167
79 tgtggcatga atactttttt taatgattga ttaaatgcaa gatagttat ctaactctgt 1227
80 gcgcaatcag ctctctatct tgacttagat tctggtggag agaagtgaga ataggcagcc 1287
81 cccaaataaa aaatatctat ggaaaaaaa aaaaaaaa 1324
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83 <211> LENGTH: 113
84 <212> TYPE: PRT
85 <213> ORGANISM: Artificial Sequence
86 <220> FEATURE:
87 <223> OTHER INFORMATION: cDNA sequence of TCL-1
88 <400> SEQUENCE: 2
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90 1 5 10 15
91 Leu Trp Ala Trp Glu Lys Phe Val Tyr Leu Asp Glu Lys Gln His Ala
92 20 25 30
93 Trp Leu Pro Leu Thr Ile Glu Ile Lys Asp Arg Leu Gln Leu Arg Val
94 35 40 45
95 Leu Leu Arg Arg Glu Asp Val Val Leu Gly Arg Pro Met Thr Pro Thr
96 50 55 60
97 Gln Ile Gly Pro Ser Leu Leu Pro Ile Met Trp Gln Leu Tyr Pro Asp
98 65 70 75 80
99 Gly Arg Tyr Arg Ser Ser Asp Ser Ser Phe Trp Arg Leu Val Tyr His
100 85 90 95
101 Ile Lys Ile Asp Gly Val Glu Asp Met Leu Leu Glu Leu Leu Pro Asp
102 100 105 110
103 Asp
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106 <212> TYPE: DNA
107 <213> ORGANISM: Artificial Sequence
108 <220> FEATURE:
109 <223> OTHER INFORMATION: genomic sequence of TCL-1
110 <400> SEQUENCE: 3
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112 ggggtcttcc agaagaagaa agggccaagg tcacccgggt gcctctccag cagcagcaga 120
113 gggcgccggt cggtgtcgct gctggccggg gcctcgagga aggcgcgggc cagctggggc 180

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123 cgccctgcgc ggaccctcac ttgccaccaa ggaccccccac aaccccgccc catccttagc 300
124 gcctgcgcgc gaccctcact tgccaccaag acccccacaa accccgcccc atcctgcctt 360
125 acgccccgcc ccaaggctgt tctcccgacc cggggctccc ccccaagacc gtcctcccg 420
126 cccgccgctt ggtggcgccc gcatgctgcc cggatataaa gggtcggccc cacatcccc 480
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139 <222> LOCATION: 108
140 <223> OTHER INFORMATION: Xaa = any amino acid
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143 <222> LOCATION: 108
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147 <222> LOCATION: 108
148 <223> OTHER INFORMATION: Xaa = Any Amino Acid
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152 1 5 10 15
153 Gln Glu Gly Ile Tyr Arg Asp Glu Tyr Gln Arg Thr Trp Val Ala Val
154 20 25 30
155 Val Glu Glu Glu Thr Ser Phe Leu Arg Ala Arg Val Gln Gln Ile Gln
156 35 40 45
157 Val Pro Leu Gly Asp Ala Ala Arg Pro Ser His Leu Leu Thr Ser Gln
158 50 55 60
159 Leu Pro Leu Met Trp Gln Leu Tyr Pro Glu Glu Arg Tyr Met Asp Asn
160 65 70 75 80
161 Asn Ser Arg Leu Trp Gln Ile Gln His His Leu Met Val Arg Gly Val
162 85 90 95
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164 100 105
167 <210> SEQ ID NO: 5
168 <211> LENGTH: 4922
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
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173 <223> OTHER INFORMATION: genomic DNA of TCL-1
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176 <222> LOCATION: (1)...(4922)
177 <223> OTHER INFORMATION: n = A,T,C or G
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W-->

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182 agcagcaagc aagaggcgccg cgggtcggtt gtcgtctgtg gccggggccc tccgaggaaa 180
183 ggcgcggrcc agctggggcc ggttctcggt tcccaggagc tgccaccgtt ccaggggagca 240
184 agtcaggcgc ggagcgttagc gcctgcgcgg gacctcact tgccaccaag rmcccccacaa 300
185 accccgcgcc atectgycct acgcccgcgc ccaaggtcgg ttctcccaga cccggggggtc 360
186 cccgccccaa ggnccgctct ccccgcctcc gccggttggt ggcgggcgca tgctgcccgg 420
187 atataaaggg tcggcccac atcccaggga ccagcgagcg gccttgagag gctctggctc 480
188 ttgcttctta ggcggcccga ggaacccatg gccagtgccc cgacactcgg ggagggcagtc 540
189 accgaccacc cggaccgcct gtgggcctgg gagaagttcg tgattttgga cgagaagcag 600
190 macgcctgcc tgcctttaac catcgaggta caaccacct tggagcggat ggcgargcag 660
191 caggggcasc cccctgggagc ttgggatncc ctagggaagg cgaggactca aggagcactc 720
192 actatggggc agggaggatc cccacagat kaagccactt ttggagcggt sctctkgagg 780
193 gatgaatagg agttctctca ggcagggaag aagggtggga aaaccccaaa ggaatttcgg 840
194 tcaaaagggg gaaccacgtg cctgtggagt gtgactataa tgttgactac agaggcatt 900
195 ttctgggctt cggggtctcta atccttaaaa atgggtatct ctaagtgact catccatag 960
196 gccgattatc ggaatcatct caggttgggt ccagaaatct gtatttttaa aaagaacccc 1020
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208 tttttcataa atcatgaaa catgcyccmc maaaaataac ccactascaaa actgtgggac 1800
209 agatttttgc tcaatcatt gaaaaggcca gcawctttt tctctcttc ttctttgtk 1860
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211 cctggcctca agcgatctcc ctgcctctgc tttccaaagc actggaatta cgtgttgag 1980
212 ccgctgcaac ccgcagaaaa aaagtgtgccc ttctatggcc ctgtctgggt ggctagacac 2040
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216 aagcggaaac tgaagtatn cctgtcctt ttctctcct nagataaagg ataggttaca 2280
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221 cyyccatccc agtssgcaa agcattwaaa aratggggga nrtwacmas tgcagatttc 2580
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228 ttatgcacaa gtggctgatc cagaagaaaa gaccacccct tctccatgaa ggcagtgcta 3000
229 accatctccc gactgtgcc atctgagggt ccttagggat gggaaggct cctgggggct 3060
230 accttaccac ccttaccac ttccggagcc aagggtcttg tagggagcc ccttttatcg

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231  cagctgctca  tatctataaa  gtacttcaca  agtttcagct  ggcactttca  ttttaccatt  3120
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233  gcaatctcag  ctaactgaaa  gctctgcctc  ccgggttcac  accattctcc  tgccctcagc  3240
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257  ccccaataaa  anaatatcca  ngggatttat  tttattnttc  cttttgggng  atnngggcag  4680
258  acattntncc  ncccgtnta  atccaatgnt  taaancccca  gtgtctcttg  aggnctacg  4740
259  tcgaanacca  ttgngtang  caacctcaaa  atttttnngt  tgnnaatnnc  cngttttcca  4800
260  gagnccccc  cntnctccat  cttnttcctn  gccnccctn  ncctcccnca  ntcccnangt  4860
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266 <212> TYPE: DNA
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
270 <223> OTHER INFORMATION: p9A primer
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277 <212> TYPE: DNA
278 <213> ORGANISM: Artificial Sequence
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Rev III primer
283 <400> SEQUENCE: 7
284 caaatggaat cctccttgcc 20
286 <210> SEQ ID NO: 8

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/14/2006
PATENT APPLICATION: US/09/441,242B TIME: 09:20:15

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:4; Xaa Pos. 108
Seq#:5; N Pos. 373,688,1155,1219,1283,1334,1335,2153,2180,2201,2224,2225
Seq#:5; N Pos. 2279,2501,2523,2545,2553,2566,2592,2599,3955,3959,3975,3976
Seq#:5; N Pos. 3982,3984,3987,3990,3992,4001,4027,4029,4072,4194,4372,4519
Seq#:5; N Pos. 4523,4584,4610,4620,4626,4632,4641,4657,4669,4673,4674,4686
Seq#:5; N Pos. 4688,4690,4691,4698,4709,4715,4734,4736,4746,4755,4759,4777
Seq#:5; N Pos. 4778,4783,4784,4789,4792,4804,4812,4814,4824,4825,4830,4835
Seq#:5; N Pos. 4840,4841,4848,4851,4856,4858,4862,4869,4890,4891,4897,4901
Seq#:5; N Pos. 4903,4906,4914
Seq#:12; Xaa Pos. 113

VERIFICATION SUMMARY

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TIME: 09:20:15

Input Set : F:\3589.1017-001 Seq List.txt

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L:30 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:33 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:138 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:142 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:146 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:150 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:96
L:175 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:179 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
L:186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:360
M:341 Repeated in SeqNo=5
L:338 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:342 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:346 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:350 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:12
L:365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:112